### Chapter 3

### Methodology

In this chapter we present our methodology to answer the research questions enunciated in ??. We investigate three research questions. In the first question, we artificially generate WebAssembly program variant, and we qualitatively compare the ability of CROW, on of the contribution of this thesis, to generate statically different, yet functionally equivalent variants. Our second research question compares the bahavior of artifically created variants through traces, execution time and preservation in machine code. The final research question evaluates the feasibility of using the program variants in security sensitive environments such as Edge-Cloud computing proposing a multivariant execution approach.

We first, enunciate the corpora of programs used to answer the research questions. We establish the metrics, and we set the configuration for the experiments. Besides, we describe the protocol for each research question.

#### 3.1 Corpora

We answer the research questions with three corpora of programs appropriate for our experiments. In Table 3.1 we listed the corpus name, the number of programs inside the corpus, the total number of functions, the range of lines of code and the original location of the corpus. In the following, we describe the filtering and description of each corpus.

1. Rosetta corpus is part of the CROW contribution []. We take programs from the Rosetta Code project<sup>1</sup>. This website hosts a curated set of solutions for specific programming tasks in various programming languages. It contains a wide range of tasks, from simple ones, such as adding two numbers, to complex algorithms like a compiler lexer. We first collect all C programs from Rosetta Code, which represents 989 programs as of 01/26/2020. We then apply a number of filters: the programs should successfully compile, they should not

<sup>1</sup>http://www.rosettacode.org/wiki/Rosetta\_Code

require user inputs, the programs should terminate and should not provide in non-deterministic results. The result of the filtering is a corpus of 303 C programs. All programs have a single function in terms of source code. These programs range from 7 to 150 lines of code and solve a variety of problems, from the Babbage problem to  $Convex\ Hull$  calculation.

- 2. **Libsodium** is part of both CROW and MEWE contributions [] []. This project is an encryption, decryption, signature and password hashing library which can be ported to WebAssembly. We selected 5 programs or endpoints to answer our research questions. These endpoints have between 8 and 2703 lines of code per function. The project is selected based on their suitability for diversity synthesis with CROW, *i.e.*, the project should have the ability to collect its modules in LLVM intermediate representation and the project should be easily portable Wasm/WASI.
- 3. **QrCode** is part of the MEWE contribution. This project is a QrCode and MicroQrCode generator written in Rust. We selected 2 programs or endpoints to answer our research questions. These endpoints have between 4 and 725 lines of code per function. As Libsodium, we select this project due to its suitability for diversity synthesis with CROW.

Corpus name	No. programs	No. functions	LOC range	Location
Rosetta	303	303	7 - 150	http:// rosettacode.org/ wiki/Rosetta_ Code
Libsodium	5	869	8 - 2703	https://github. com/jedisct1/ libsodium
QrCode	2	1849	4 - 725	https://github. com/kennytm/ qrcode-rust
Total	310	3021		

Table 3.1: Corpora description. The table is composed by the name of the corpus, the number of programs, the number of functions, the lines of code range and the location of the corpus.

## 3.2 RQ1. To what extent we can generate program variants for WebAssembly.

In this research question, we investigate whether we can artifically generate program variants for WebAssembly. We use CROW to generate program variants from an

original program, written in C/C++ or directly passing a LLVM bitcode module to it. We analyze the properties the programs should have to generate a handful number of variants.

#### 3.2.1 Metrics

To assess our approach's ability to generate WebAssembly binaries that are statically different, we compute the number of unique variants generated by CROW for each original function. For each program and its variants we calculate the count of unique generated variants.

**Metric 1** Population size S(P): Given a program P and its generated variants V, the population size metric is defined as.

$$S(P, V) = |V \cap \{P\}|$$

#### 3.2.2 Protocol

The generation of program variants is part of the answering of all our research questions. Therefore, the first step and research question is related to the ability of CROW to generate a handful number of program variants.

We pass each of the 303+869+1849 functions in the corpora to CROW, which synthesizes program variants with an enumerative strategy. All possible programs that can be generated for a given language (LLVM in the case) are constructed and verified for equivalence. There are two parameters to control the size of the search space and hence the time required to traverse it. On the one hand, one can limit the size of the variants. On the other hand, one can limit the set of instructions used for the synthesis. On the other hand, in our experiments, we use between 1 instruction (only additions) and 60 instructions (all supported instructions in the synthesizer).

These two configuration parameters allow the user to find a trade-off between the number of variants that are synthesized and the time taken to produce them. In Table 3.2 we listed the configuration for both corpora. For the current evaluation, given the size of the corpus, we set the exploration time to 1 hour maximum per function for Rosetta . In the cases of Libsodium and QrCode, we set the timeout to 5 minutes per function in the exploration stage. We set all 60 supported instructions in CROW for Rosetta , Libsodium and QrCode corpora.

CORPUS Name	Exploration timeout	Max. instructions
Rosetta	1h	60
Libsodium	$5\mathrm{m}$	60
$\operatorname{QrCode}$	5m	60

Table 3.2: CROW tweaking for variants generation. The table is composed by the name of the corpus, the timeout parameter and the count of allowed instructions during the synthesis process.

We pass each function of Table 3.1 corpora to CROW, and we evaluate the Metric 1 for each of them in order to answer RQ1.

### 3.3 RQ2. To what extent the artifically generated variants are different?

In this second research question, we investigate to what extent the artificially created variants are different between them and to the original program. To conduct this research question, we perform three experiments: static comparison, dynamic comparison and variant's preservation. We use the original programs from corpora described in section 3.1 and their variants generated in the answering of RQ1.

#### 3.3.1 Metrics

A program and its variants compose what we call a program's population. Notice that all metric that can be proposed over the programs generated in RQ1 make sense only at the population level. It only makes sense to compare programs that are in fact semantically equivalent, i.e. from the same population.

To compare two programs of the same population we propose a global alignment approach using Dynamic Time Warping (DTW). In a previous work of us, we highlighted how this approach can be used to measure similarity between program traces  $\cite{Plance}$ . Dynamic Time Warping  $\cite{Plance}$  computes the global alignment between two sequences. It returns a value capturing the cost of this alignment, which is actually a distance metric. The larger the DTW distance, the more different the two sequences are. In the following we define the  $dt\_static$  metric.

**Metric 2** dt\_static: Given two programs of the same program's population  $P_X$  and  $V_X$  written in X code, dt\_static( $P_X$ ,  $V_X$ ), computes the DTW distance between the corresponding program instructions for representation X.

A  $dt\_static(P_X, V_X)$  of 0 means that the code of both the original program and the variant is the same, i.e., they are statically identical in the representation X.

The higher the value of dt\_static, the more different the programs are in representation X.

Notice that, for comparing WebAssembly programs, the metric is the instantiation of  $dt\_static$  with X = WebAssembly.

We measure the difference between programs at runtime by comparing their execution traces also with and alignment metric, at function and instruction level. Besides, we collect the execution times of WebAssembly programs.

**Metric 3**  $dt\_dyn$ : Given a program P, a CROW generated variant P' and T a trace space  $(T \in \{Function, Instruction\})$   $dt\_dyn(P,P',T)$ , computes the DTW distance between the traces collected during their execution in the T space. A  $dt\_dyn$  of 0 means that both traces are identical.

The higher the value, the more different the traces.

**Metric 4** Execution time: Given a WebAssembly program P, the execution time is the time spent to execute the binary.

WebAssembly is an intermediate language and interpreters produce machine code to actually execute them. For program variants, this means that compiling can undo artificial introduced transformations, for example, through optimization passes. When a code transformation for a variant is maintained from the first time it is introduced to the final machine code generation is then a preserved variant. For this mentioned reasoning we need a preservation metric. The key property we consider is as follows:

**Property 1** Preservation: Given a program P and a variant V' from the same program's population, if  $dt\_static(P_{Wasm}, P'_{Wasm}) > 0$  and  $dt\_static(P_{x86}, P'_{x86}) > 0 \implies both programs are still different when compiled to machine code.$ 

If the property fits for two programs, then the underlying compiler does not remove the transformations made by CROW.

Metric 5 Preservation: Given a WebAssembly programs P and a collection of generated variants V, the preservation ratio is the number of pair of programs that fit with Property 1 for a specific compiling engine over the total number of program pairs.

#### 3.3.2 Protocol

For each program's population generated in the answering of RQ1, we compare the sequence of instructions of each variant with the initial program and the other variants. We obtain the Metric 2 values for each program-variant WebAssembly pair code. We compute the DTW distances with STRAC [?].

To compare program and variants behavior during runtime, we analyze all the unique program variants generated by CROW in a pairwise comparison as well. We use SWAM<sup>2</sup> to execute each program and variant and to collect the function and instruction traces. SWAM is a WebAssembly interpreter that provides functionalities to capture the dynamic information of WebAssembly program executions including the stack operations. We compute the DTW distances with STRAC [?].

Furthermore, we collect the execution time, Metric 4, for all programs and their variants. We execute each program or variant 10000 times, and we compare the collected execution time distributions using a Mann-Withney test [?] in a pairwise strategy.

We collect Metric 5 for all program pairs in all program's populations. We use two WebAssembly engines to study variant's preservation after compilling.

- V8 [?]: the engine used by Chrome and NodeJS to execute JavaScript and WebAssembly.
- wasmtime [?]: a standalone runtime for WebAssembly. This engine is used by the Fastly platform to provide Edge-Cloud computing services.

We only take into account the x86 representation after the WebAssembly code is compiled to the machine code. This decision is not arbitrary, according to the study of TODO Paper on binary diff survey, any conclusion carried out by comparing two program binaries under a specific target can be extrapolated to another target for the same binaries.

### 3.4 RQ3. To what extent the artificially generated variants can be used to enforce security?

In the last research question we study whether the created variants can be used in real-world applications and what properties offer the composition of the variants as multivariant execution binaries. For this purpose, we build multivariant binaries to be deployed at the Edge in the Fastly platform. We use the variants generated for the programs of the Libsodium and QrCode corpora, 2+5 programs involving 869+1849 functions respectively. Multivariant binaries are created by converting each program's population into a single function for which each call at runtime selects and executes a different variant. One of the contributions of this work is MEWE, a tool that automatically creates multivariant binaries out of program variants generated by CROW.

With this research question, we also assess the ability of MEWE, to produce binaries that actually exhibit random execution paths when executed on one edge node. We check the diversity of execution traces gathered from the execution of a multivariant binary. The traces are collected from all edge nodes in order to assess

<sup>&</sup>lt;sup>2</sup>https://github.com/satabin/swam

Multivariant Execution (MVE) at a worldwide scale. MEWE generates binaries that embed a multivariant behavior. We measure to what extent MEWE generates different execution times on the edge. Then, we discuss how multivariant binaries contribute to less predictable timing side-channels.

#### 3.4.1 Metrics

In the following we enunciate the metric use to answer our last research question. To compare the diversity of function traces for the 7 created multivariant binaries we enunciate the following metrics.

**Metric 6** Unique traces: R(n, e). Let  $S(n, e) = \{T_1, T_2, ..., T_{100}\}$  be the collection of 100 traces collected for one endpoint e on an edge node n, H(n, e) the collection of hashes of each trace and U(n, e) the set of unique trace hashes in H(n, e). The uniqueness ratio of traces collected for edge node n and endpoint e is defined as

$$R(n,e) = \frac{|U(n,e)|}{|H(n,e)|}$$

**Metric 7** Normalized Shannon entropy: E(e) Let e be an endpoint,  $C(e) = {}^{64}_{n=0}H(n,e)$  be the union of all trace hashes for all edge nodes. The normalized Shannon Entropy for the endpoint e over the collected traces is defined as:

$$E(e) = -\sum \frac{p_x * log(p_x)}{log(|C(e)|)}$$

Where  $p_x$  is the discrete probability of the occurrence of the hash x over C(e).

Notice that we normalize the standard definition of the Shannon Entropy, Metric 7, by using the perfect case where all trace hashes are different. This normalization allows us to compare the calculated entropy between endpoints. The value of the metric can go from 0 to 1. The worst entropy, value 0, means that the endpoint always perform the same path independently of the edge node and the number of times the trace is collected for the same node. On the contrary, 1 for the best entropy, when each edge node executes a different path every time the endpoint is requested.

#### 3.4.2 Protocol

We run the experiments to answer RQ3 on the Fastly edge computing platform. We deploy and execute the original and the multivariant endpoints on 64 edge nodes located around the world<sup>3</sup>. These edge nodes usually have an arbitrary and heterogeneous composition in terms of architecture and CPU model.

<sup>&</sup>lt;sup>3</sup>The number of nodes provided in the whole platform is 72, we decided to keep only the 64 nodes that remained stable during our experimentation.

11

We execute each endpoint, multiple times on each node, to measure the diversity of execution traces that are exhibited by the multivariant binaries. Each query on the same endpoint is performed with the same input value. This is to guarantee that, if we observe different traces for different executions, it is due to the presence of multiple function variants. The inputs that we pass to execute the endpoints at the edge and the received output for all executions are available in the reproduction repository at TODO REPO.

For each query, we collect the execution trace, i.e., the sequence of function names that have been executed when triggering the query. To observe these traces, we instrument the multivariant binaries to record each function entrance.

We then measure the number of unique execution traces exhibited by each multivariant binary, Metric 6, on each separate edge node. To compare the traces, we hash them with the md5sum function. We then calculate the number of unique hashes among the 100 traces collected for an endpoint on one edge node. We follow by collecting the normalized Shannon entropy, Metric 7, for all collected execution traces for each endpoint. The Shannon Entropy gives the uncertainty in the outcome of a sampling process. If a specific trace has a high frequency of appearing in part of the sampling, then it is certain that this trace will appear in the other part of the sampling.

We calculate Metric 7 for the 7 endpoints, for 100 traces collected from 64 edge nodes, for a total of 6400 collected traces per endpoint. Each trace is collected in a round robin strategy, i.e., the traces are collected from the 64 edge nodes sequentially. For example, we collect the first trace from all nodes before continuing to the collection of the second trace. This process is followed until 100 traces are collected from all edge nodes.

In addition, we collect 100k execution times for each binary, both the original and multivariant binaries. We perform a Mann-Withney U test [?] to compare both execution time distributions. If the P-value is lower than 0.05, two compared distributions are different.

#### 3.5 Conclusions

In this chapter we present the methodology we follow to answer our three research questions. We first propose to measure the ability of CROW to generate variants out of 3021 functions. Then, we propose to collect and reuse the generateed variants to study to what extent they offer different observable behavior through static, dynamic and variant's preservation. Finally, we propose a protocol to study the impact of composition of the generated variants in a single binary deployed at the Edge. In this study, we enumerate and enunciate the properties and metrics that might lead us to answer the impact of automatic diversification for WebAssembly programs. In the next chapter, we present and discuss the results obtained with this experimental protocol.

# Appended papers